



## **Sample Collection, Handling and Analysis of *Escherichia Coli***

Montana Department of Environmental Quality

Water Quality Planning Section

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## Escherichia coli

This Standard Operating Procedure (SOP) describes methods developed for measuring Escherichia coli (*E. coli*) levels for use in water quality assessments. Before using these methods, please inquire about any recent SOP revisions by contacting the following individuals (or their replacements):

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### ***1. Scope and Applicability***

The methods described herein are used in any waterbody or source of drinking water.

#### **1.1. General Considerations**

Before conducting a water quality survey, determine specific data quality objectives (DQOs) and clearly define the information sought. DQOs are qualitative and quantitative statements developed to specify the quality of data needed to support specific decisions and conclusions about the information sought.

#### **1.2. Site Selection**

The total number of sites, their location in the study area, and the number of replicate samples used to represent each site should be determined as part of the sample process design during the DQO process. Once established, the design should be adhered to in order to achieve the objectives of the project. In developing a sampling design, always consider potential constraints to collecting representative samples such as site access. Document any changes in study design due to site access restrictions.

The study design should specify requisite site specifications for the study. For example, many study designs use a beach area or fishing access point as the specific point of comparison between streams. A rationale described in the Sampling and Analysis Plan (SAP) for determining representative sampling areas for the anticipated stream types would be a site selection specification.

### ***2. Sample Collection Protocols***

#### **2.1. Surface Waters**

The species *Escherichia coli* (*E. coli*) is the principal indicator of suitability of a waterbody for recreational use.

*E. coli* are analyzed at the Public Health Laboratory or in the field using the IDEXX, Inc. Colilert™ method<sup>1</sup>. Coliform densities are reported as most probable number count per 100 ml. This method must be performed by qualified state employees or employees trained by qualified employees.

### **2.1.1 Equipment**

1. IDEXX Collection bottles for coliform bacteria (100ml). Sterilized and treated with sodium thiosulfate. These bottles are available at the Public Health Laboratory or from IDEXX. DEQ has many of these bottles in the Water Laboratory.



2. Colilert™ packets: You will need one packet per sample



3. Quanti-Tray® /2000 sample trays: You will need one Quanti-Tray per sample.



4. Quanti-Tray® Sealer. If you are performing the Colilert method in the field, you will need a source of electricity for the sealer.



5. Quanti-Tray® /2000 rubber insert for the sealer.



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<sup>1</sup> Note: There are many different methods that evaluate *E. coli* levels in surface water. Use any *E. coli* method listed in the most recent version of the 40 CFR Part 136.

6. Incubator (Millipore™ Dual Chamber Portable Incubator): The portable incubator can be powered via either a wall outlet, car battery or an internal battery. The internal battery will only last six hours.



7. Cooler and ice
8. Portable battery powered black light



9. Scissors
10. Heavy Blanket
11. Sharpie marker
12. Autoclaved 250 ml container

### **2.1.2 Sample Collection**

1. Label the bottle using the sharpie with the STORET Station ID and replicate number.
2. Remove the tamper seal from the bottle lid.
3. Do not rinse the sterile Colilert sample bottle!
4. Use a satellite (autoclaved) 250 ml container to physically collect the sample from the waterbody. Fill the satellite container to 200 ml, cap the container, and shake vigorously.
5. Fill the sterile bottle to the 100 ml line from the satellite container. For splits, fill two Colilert containers.
6. Enter required data onto the DEQ Site Visit Form.
7. Place the sample jar in a cooler with ice or refrigerate until you process the sample.

### **2.1.3 Sample Processing**

All steps to the Colilert™ method are listed on the IDEXX, Inc. Website:

<http://www.idexx.com/water/colilert/index.jsp>

1. Add the Colilert™ reagent to the sample close the lid tightly and shake well.



2. Pour the sample mixed with the reagent into Quanti-Tray®/2000.



3. Place the full Quanti-Tray/2000 in the rubber insert into the Sealer.



4. After the sealer has completed the cycle, inspect the sealed Quanti-Tray/2000 to make sure it was completely sealed.
5. Cut the sealed Quanti-Tray/2000 in half with the scissors. Be sure to cut in between the row of cells. Do not damage or cut any of the cells.
6. Place the cut Quanti-Tray/2000 in the incubator at  $35 \pm 0.5$  degrees Celsius for 24 to 28 hours.
7. Read results from Quanti-Tray/2000:
  - a. The yellow cells indicate that coliform bacteria are present. Count the number of large and small yellow cells on the Quanti-Tray/2000. Use the Most Probable Number (MPN) tables in Appendix A to determine the MPN count per 100 ml of sample. Record all information on the sample collection form (Appendix B).



- b. The yellow cells with *E. coli* present will fluoresce under a black light. When you count the cells in the field, cover the Quanti-Tray/2000 and your head with the heavy blanket and count the number of yellow cells that fluoresce when exposed to the black light. Use the MPN tables in Appendix A to determine the MPN count per 100 ml of sample. Record all information on the sample collection form (Appendix B).

#### **2.1.4 Sample Preservation**

1. If the samples will not be processed immediately, bacteria samples should be iced or refrigerated at a temperature of < 10 degrees Celsius (APHA 2005)
2. *E. coli* samples must not be held more than six hours between collection and initiation of analyses. Samples must be processed within 2 hours of arriving to a lab (APHA 2005).

#### **2.1.5 Precautions**

1. When you sample surface waters, try to collect the water from a depth of 15.2 cm (6 in), while still leaving room at the top of the bottle for mixing. Avoid surface scum. Do not touch the insides of the bottle; keep the hands near the base of the bottle while sampling. Tightly close the lid and shake the bottle to homogenize the sample.
2. If any cells leak after you cut the Quanti-Tray, you must restart the entire process. Therefore, collect more than one sample at a time just in case you need to start over.

#### **2.1.6 Quality Control**

Examples of quality control for *E. coli* samples are as follows:

##### **2.1.6.1 Field Splits**

Field splits are collected to measure total method error and should always be included in the sampling design at a predetermined frequency (usually 30%).

Field splits are two or more samples separated from a larger sample. Split samples are processed in the field or laboratory as discrete samples and their difference evaluated. This difference represents total method error (e.g., heterogeneity of site, reproducibility of sampling technique, sorting error, and identification error).

Relative Percent Difference (RPD) is used to evaluate results differences between two replicate samples.



$$RPD = \frac{|(result1 - result2)|}{(result1 + result2)/2} \times 100$$

Note: Bacteria replicates are typically variable in nature (Wymer et al. 2005). Use the RPD calculation only as a descriptive statistic for bacteria data.

### **2.1.7 Data Analysis and Interpretation**

In order to determine if *E. coli* levels for a waterbody exceed the water quality standard, you have to compare the sample results against two benchmarks. The first is the value that the geometric mean of multiple samples cannot exceed. The other is the maximum value that 10 percent of the samples cannot exceed.

#### **2.1.7.1 Calculating the Geometric Mean**

The geometric mean is based on a minimum of five samples obtained during separate 24-hour periods during any consecutive 30-day period analyzed by the most probable number or equivalent membrane filter methods (ARM<sup>2</sup> 17.30.620.2).

The geometric mean is the value obtained by taking the Nth root of the product of the measured values where zero values for measured values are taken to be the detection limit (BBNEP 2006; ARM 17.30.602.13).

$$\sqrt[n]{\prod_{i=1}^n X_i}$$

#### **2.1.7.2 Data Interpretation**

Use the following steps to determine if a waterbody meets its contact recreation use:

1. Find out which classification is assigned to your waterbody in ARM 17.30.607 - 17.30.615 (e.g. A-1, B-1, etc.).
2. View the standards listed in Tables 1 and 2 and find the appropriate water use classification
3. Determine if the waterbody meets its assigned contact recreation use
  - a. Determine when the data were collected:
    - i. If the data were collected between April 1 to October 31, use the *E. coli* standards listed Table 1 to determine if the waterbody is meeting its contact recreation use.
    - ii. If the data were collected from November 1 through March 31, use the *E. coli* standards listed in Table 2 to determine if the waterbody is meeting its contact recreation use. If you have five or more data points, summarize them into a geometric mean and compare it to

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<sup>2</sup> ARM = Administrative Rules of Montana

the *E. coli* “geometric mean” standard. Then, compare each individual data point to the “10% level”. If either the “geometric mean” or the “10%” standards are exceeded, then the waterbody is not meeting its contact recreation use.

Table 1. Use this table to interpret *E. coli* data collected between April 1 and October 31.

	<i>E. coli</i> (10% of all samples must not exceed this <i>E. coli</i> (geometric mean; density, CFU*/100ml) CFU*/100ml)	
Contact Recreation Standard		
<b>A-closed; A-1</b>	32	64
<b>B-1; B-2; B-3; C-1; C-2; I; C-3</b>	126	252

\* CFU = Colony-Forming-Units

Table 2. Use this table to interpret *E. coli* data collected between November 1 and March 31.

	<i>E. coli</i> (10% of all samples must not exceed this <i>E. coli</i> (geometric mean; density, CFU*/100 ml) CFU*/100ml)	
Contact Recreation Standard		
<b>A-closed; A-1</b>	32	64
<b>B-1; B-2; B-3; C-1; C-2; I; C-3</b>	630	1260

\* CFU = Colony-Forming-Units

- iii. If you have one or more data points, compare each individual data point to the “10% level”. If 10% of the values are greater than the 10% level, then the waterbody is not meeting its contact recreation use.

## Literature Cited

- American Public Health Association (APHA). 2005. Standard methods for the examination of water and wastewater. 21st ed. American Public Health Association, American Water Works Association, and Water Pollution Control Federation. Washington, DC.
- Buzzards Bay National Estuary Program (BBNEP). 2006. Calculating Geometric Means. Website: <http://www.buzzardsbay.org/geomean.htm> .
- Wymer, L.J., K.P. Brenner, J.W. Martinson, W.R. Stutts, S.A. Schaub and A. P. Dufour. 2005. The EMPACT Beaches Project: Results from a Study on Microbiological Monitoring in Recreational Waters. United States Environmental Protection Agency, Office of Research and Development, National Exposure Research Laboratory, Cincinnati, OH USA. EPA 600/R-04/023.

## Appendix A: The Most Probable Number Table for the Colilert method

# Large Wells Positive	IDEXX Quanti-Tray®/2000 MPN Table																								
Wells Positive	# Small Wells Positive																								
	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24
0	<1	1.0	2.0	3.0	4.0	5.0	6.0	7.0	8.0	9.0	10.0	11.0	12.0	13.0	14.1	15.1	16.1	17.1	18.1	19.1	20.2	21.2	22.2	23.3	24.3
1	1.0	2.0	3.0	4.0	5.0	6.0	7.1	8.1	9.1	10.1	11.1	12.1	13.2	14.2	15.2	16.2	17.3	18.3	19.3	20.4	21.4	22.4	23.5	24.5	25.6
2	2.0	3.0	4.1	5.1	6.1	7.1	8.1	9.2	10.2	11.2	12.2	13.3	14.3	15.4	16.4	17.4	18.5	19.5	20.6	21.6	22.7	23.7	24.8	25.8	26.9
3	3.1	4.1	5.1	6.1	7.2	8.2	9.2	10.3	11.3	12.4	13.4	14.5	15.5	16.5	17.6	18.6	19.7	20.8	21.8	22.9	23.9	25.0	26.1	27.1	28.2
4	4.1	5.2	6.2	7.2	8.3	9.3	10.4	11.4	12.5	13.5	14.6	15.6	16.7	17.8	18.8	19.9	21.0	22.0	23.1	24.2	25.3	26.3	27.4	28.5	29.6
5	5.2	6.3	7.3	8.4	9.4	10.5	11.5	12.6	13.7	14.7	15.8	16.9	17.9	19.0	20.1	21.2	22.2	23.3	24.4	25.5	26.6	27.7	28.8	29.9	31.0
6	6.3	7.4	8.4	9.5	10.6	11.6	12.7	13.8	14.9	16.0	17.0	18.1	19.2	20.3	21.4	22.5	23.6	24.7	25.8	26.9	28.0	29.1	30.2	31.3	32.4
7	7.5	8.5	9.6	10.7	11.8	12.8	13.9	15.0	16.1	17.2	18.3	19.4	20.5	21.6	22.7	23.8	24.9	26.0	27.1	28.3	29.4	30.5	31.6	32.8	33.9
8	8.6	9.7	10.8	11.9	13.0	14.1	15.2	16.3	17.4	18.5	19.6	20.7	21.8	22.9	24.1	25.2	26.3	27.4	28.6	29.7	30.8	32.0	33.1	34.3	35.4
9	9.8	10.9	12.0	13.1	14.2	15.3	16.4	17.6	18.7	19.8	20.9	22.0	23.2	24.3	25.4	26.5	27.7	28.9	30.0	31.2	32.3	33.5	34.6	35.8	37.0
10	11.0	12.1	13.2	14.4	15.5	16.6	17.7	18.9	20.0	21.1	22.3	23.4	24.6	25.7	26.9	28.0	29.2	30.3	31.5	32.7	33.8	35.0	36.2	37.4	38.6
11	12.2	13.4	14.5	15.6	16.8	17.9	19.1	20.2	21.4	22.5	23.7	24.8	26.0	27.2	28.3	29.5	30.7	31.9	33.0	34.2	35.4	36.6	37.8	39.0	40.2
12	13.5	14.6	15.8	16.9	18.1	19.3	20.4	21.6	22.8	23.9	25.1	26.3	27.5	28.6	29.8	31.0	32.2	33.4	34.6	35.8	37.0	38.2	39.5	40.7	41.9
13	14.8	16.0	17.1	18.3	19.5	20.6	21.8	23.0	24.2	25.4	26.6	27.8	29.0	30.2	31.4	32.6	33.8	35.0	36.2	37.5	38.7	39.9	41.2	42.4	43.6
14	16.1	17.3	18.5	19.7	20.9	22.1	23.3	24.5	25.7	26.9	28.1	29.3	30.5	31.7	33.0	34.2	35.4	36.7	37.9	39.1	40.4	41.6	42.9	44.2	45.4
15	17.5	18.7	19.9	21.1	22.3	23.5	24.7	25.9	27.2	28.4	29.6	30.9	32.1	33.3	34.6	35.8	37.1	38.4	39.6	40.9	42.2	43.4	44.7	46.0	47.3
16	18.9	20.1	21.3	22.6	23.8	25.0	26.2	27.5	28.7	30.0	31.2	32.5	33.7	35.0	36.3	37.5	38.8	40.1	41.4	42.7	44.0	45.3	46.6	47.9	49.2
17	20.3	21.6	22.8	24.1	25.3	26.6	27.8	29.1	30.3	31.6	32.9	34.1	35.4	36.7	38.0	39.3	40.6	41.9	43.2	44.5	45.9	47.2	48.5	49.8	51.2
18	21.8	23.1	24.3	25.6	26.9	28.1	29.4	30.7	32.0	33.3	34.6	35.9	37.2	38.5	39.8	41.1	42.4	43.8	45.1	46.5	47.8	49.2	50.5	51.9	53.2
19	23.3	24.6	25.9	27.2	28.5	29.8	31.1	32.4	33.7	35.0	36.3	37.6	39.0	40.3	41.6	43.0	44.3	45.7	47.1	48.4	49.8	51.2	52.6	54.0	55.4
20	24.9	26.2	27.5	28.8	30.1	31.5	32.8	34.1	35.4	36.8	38.1	39.5	40.8	42.2	43.6	44.9	46.3	47.7	49.1	50.5	51.9	53.3	54.7	56.1	57.6
21	26.5	27.9	29.2	30.5	31.8	33.2	34.5	35.9	37.3	38.6	40.0	41.4	42.8	44.1	45.5	46.9	48.4	49.8	51.2	52.6	54.1	55.5	56.9	58.4	59.9
22	28.2	29.5	30.9	32.3	33.6	35.0	36.4	37.7	39.1	40.5	41.9	43.3	44.8	46.2	47.6	49.0	50.5	51.9	53.4	54.8	56.3	57.8	59.3	60.8	62.3
23	29.9	31.3	32.7	34.1	35.5	36.8	38.3	39.7	41.1	42.5	43.9	45.4	46.8	48.3	49.7	51.2	52.7	54.2	55.6	57.1	58.6	60.2	61.7	63.2	64.7
24	31.7	33.1	34.5	35.9	37.3	38.8	40.2	41.7	43.1	44.6	46.0	47.5	49.0	50.5	52.0	53.5	55.0	56.5	58.0	59.5	61.1	62.6	64.2	65.8	67.3
25	33.6	35.0	36.4	37.9	39.3	40.8	42.2	43.7	45.2	46.7	48.2	49.7	51.2	52.7	54.3	55.8	57.3	58.9	60.5	62.0	63.6	65.2	66.8	68.4	70.0
26	35.5	36.9	38.4	39.9	41.4	42.8	44.3	45.9	47.4	48.9	50.4	52.0	53.5	55.1	56.7	58.2	59.8	61.4	63.0	64.7	66.3	67.9	69.6	71.2	72.9
27	37.4	38.9	40.4	42.0	43.5	45.0	46.5	48.1	49.6	51.2	52.8	54.4	56.0	57.6	59.2	60.8	62.4	64.1	65.7	67.4	69.1	70.8	72.5	74.2	75.9
28	39.5	41.0	42.6	44.1	45.7	47.3	48.8	50.4	52.0	53.6	55.2	56.9	58.5	60.2	61.8	63.5	65.2	66.9	68.6	70.3	72.0	73.7	75.5	77.3	79.0
29	41.7	43.2	44.8	46.4	48.0	49.6	51.2	52.8	54.5	56.1	57.8	59.5	61.2	62.9	64.6	66.3	68.0	69.8	71.5	73.3	75.1	76.9	78.7	80.5	82.4
30	43.9	45.5	47.1	48.7	50.4	52.0	53.7	55.4	57.1	58.8	60.5	62.2	64.0	65.7	67.5	69.3	71.0	72.9	74.7	76.5	78.3	80.2	82.1	84.0	85.9
31	46.2	47.9	49.5	51.2	52.9	54.6	56.3	58.1	59.8	61.6	63.3	65.1	66.9	68.7	70.5	72.4	74.2	76.1	78.0	79.9	81.8	83.7	85.7	87.6	89.6
32	48.7	50.4	52.1	53.8	55.6	57.3	59.1	60.9	62.7	64.5	66.3	68.2	70.0	71.9	73.8	75.7	77.6	79.5	81.5	83.5	85.4	87.5	89.5	91.5	93.6
33	51.2	53.0	54.8	56.5	58.3	60.2	62.0	63.8	65.7	67.6	69.5	71.4	73.3	75.2	77.2	79.2	81.2	83.2	85.2	87.3	89.3	91.4	93.5	95.7	97.8
34	53.9	55.7	57.6	59.4	61.3	63.1	65.0	67.0	68.9	70.8	72.8	74.8	76.8	78.8	80.8	82.9	85.0	87.1	89.2	91.4	93.5	95.7	97.9	100.2	102.4
35	56.8	58.6	60.5	62.4	64.4	66.3	68.3	70.3	72.3	74.3	76.3	78.4	80.5	82.6	84.7	86.9	89.1	91.3	93.5	95.7	98.0	100.3	102.6	105.0	107.3
36	59.8	61.7	63.7	65.7	67.7	69.7	71.7	73.8	75.9	78.0	80.1	82.3	84.5	86.7	88.9	91.2	93.5	95.8	98.1	100.5	102.9	105.3	107.7	110.2	112.7
37	62.9	65.0	67.0	69.1	71.2	73.3	75.4	77.6	79.8	82.0	84.2	86.5	88.8	91.1	93.4	95.8	98.2	100.6	103.1	105.6	108.1	110.7	113.3	115.9	118.6
38	66.3	68.4	70.6	72.7	74.9	77.1	79.4	81.6	83.9	86.2	88.6	91.0	93.4	95.8	98.3	100.8	103.4	105.9	108.6	111.2	113.9	116.6	119.4	122.2	125.0
39	70.0	72.2	74.4	76.7	78.9	81.3	83.6	86.0	88.4	90.9	93.4	95.9	98.4	101.0	103.6	106.3	109.0	111.8	114.6	117.4	120.3	123.2	126.1	129.2	132.2
40	73.8	76.2	78.6	80.9	83.3	85.7	88.2	90.6	93.3	95.9	98.5	101.2	103.9	106.7	109.5	112.4	115.3	118.2	121.2	124.3	127.4	130.5	133.7	137.0	140.3
41	78.0	80.5	83.0	85.5	88.0	90.6	93.3	95.9	98.7	101.4	104.3	107.1	110.0	113.0	116.0	119.1	122.2	125.4	128.7	132.0	135.4	138.8	142.3	145.9	149.5
42	82.6	85.2	87.8	90.5	93.2	96.0	98.8	101.7	104.6	107.6	110.6	113.7	116.9	120.1	123.4	126.7	130.1	133.6	137.2	140.8	144.5	148.3	152.2	156.1	160.2
43	87.6	90.4	93.2	96.0	99.0	101.9	105.0	108.1	111.2	114.5	117.8	121.1	124.6	128.1	131.7	135.4	139.1	143.0	147.0	151.0	155.2	159.4	163.8	168.2	172.8
44	93.1	96.1	99.1	102.2	105.4	108.6	111.9	115.3	118.7	122.3	125.9	129.6	133.4	137.4	141.4	145.5	149.7	154.1	158.5	163.1	167.9				

# Large Wells Positive	IDEXX Quanti-Tray®/2000 MPN Table																							
	# Small Wells Positive																							
	26	28	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48
0	25.3	26.4	27.4	28.4	29.5	30.5	31.5	32.6	33.6	34.7	35.7	36.8	37.8	38.9	40.0	41.0	42.1	43.1	44.2	45.3	46.3	47.4	48.5	49.5
1	26.6	27.7	28.7	29.8	30.8	31.9	32.9	34.0	35.0	36.1	37.2	38.2	39.3	40.4	41.4	42.5	43.6	44.7	45.7	46.8	47.9	49.0	50.1	51.2
2	27.9	29.0	30.0	31.1	32.2	33.2	34.3	35.4	36.5	37.5	38.6	39.7	40.8	41.9	43.0	44.0	45.1	46.2	47.3	48.4	49.5	50.6	51.7	52.8
3	29.3	30.4	31.4	32.5	33.6	34.7	35.8	36.8	37.9	39.0	40.1	41.2	42.3	43.4	44.5	45.6	46.7	47.8	48.9	50.0	51.2	52.3	53.4	54.5
4	30.7	31.8	32.8	33.9	35.0	36.1	37.2	38.3	39.4	40.5	41.6	42.8	43.9	45.0	46.1	47.2	48.3	49.5	50.6	51.7	52.9	54.0	55.1	56.3
5	32.1	33.2	34.3	35.4	36.5	37.6	38.7	39.9	41.0	42.1	43.2	44.4	45.5	46.6	47.7	48.9	50.0	51.2	52.3	53.5	54.6	55.8	56.9	58.1
6	33.5	34.7	35.8	36.9	38.0	39.2	40.3	41.4	42.6	43.7	44.8	46.0	47.1	48.3	49.4	50.6	51.7	52.9	54.1	55.2	56.4	57.6	58.7	59.9
7	35.0	36.2	37.3	38.4	39.6	40.7	41.9	43.0	44.2	45.3	46.5	47.7	48.8	50.0	51.2	52.3	53.5	54.7	55.9	57.1	58.3	59.4	60.6	61.8
8	36.6	37.7	38.9	40.0	41.2	42.3	43.5	44.7	45.9	47.0	48.2	49.4	50.6	51.8	53.0	54.1	55.3	56.5	57.7	59.0	60.2	61.4	62.6	63.8
9	38.1	39.3	40.5	41.6	42.8	44.0	45.2	46.4	47.6	48.8	50.0	51.2	52.4	53.6	54.8	56.0	57.2	58.4	59.7	60.9	62.1	63.4	64.6	65.8
10	39.7	40.9	42.1	43.3	44.5	45.7	46.9	48.1	49.3	50.6	51.8	53.0	54.2	55.5	56.7	57.9	59.2	60.4	61.7	62.9	64.2	65.4	66.7	67.9
11	41.4	42.6	43.8	45.0	46.3	47.5	48.7	49.9	51.2	52.4	53.7	54.9	56.1	57.4	58.6	59.9	61.2	62.4	63.7	65.0	66.3	67.5	68.8	70.1
12	43.1	44.3	45.6	46.8	48.1	49.3	50.6	51.8	53.1	54.3	55.6	56.8	58.1	59.4	60.7	62.0	63.2	64.5	65.8	67.1	68.4	69.7	71.0	72.4
13	44.9	46.1	47.4	48.6	49.9	51.2	52.5	53.7	55.0	56.3	57.6	58.9	60.2	61.5	62.8	64.1	65.4	66.7	68.0	69.3	70.7	72.0	73.3	74.7
14	46.7	48.0	49.3	50.5	51.8	53.1	54.4	55.7	57.0	58.3	59.6	60.9	62.3	63.6	64.9	66.3	67.6	68.9	70.3	71.6	73.0	74.4	75.7	77.1
15	48.6	49.9	51.2	52.5	53.8	55.1	56.4	57.8	59.1	60.4	61.8	63.1	64.5	65.8	67.2	68.6	69.9	71.3	72.6	74.0	75.4	76.8	78.2	79.6
16	50.5	51.8	53.2	54.5	55.8	57.2	58.5	59.9	61.2	62.6	64.0	65.3	66.7	68.1	69.5	70.9	72.3	73.7	75.1	76.5	77.9	79.3	80.8	82.2
17	52.5	53.9	55.2	56.6	58.0	59.3	60.7	62.1	63.5	64.9	66.3	67.7	69.1	70.5	71.9	73.3	74.8	76.2	77.6	79.1	80.5	82.0	83.5	84.9
18	54.6	56.0	57.4	58.8	60.2	61.6	63.0	64.4	65.8	67.2	68.6	70.1	71.5	73.0	74.4	75.9	77.3	78.8	80.3	81.8	83.3	84.8	86.3	87.8
19	56.8	58.2	59.6	61.0	62.4	63.9	65.3	66.8	68.2	69.7	71.1	72.6	74.1	75.5	77.0	78.5	80.0	81.5	83.1	84.6	86.1	87.6	89.2	90.7
20	59.0	60.4	61.9	63.3	64.8	66.3	67.7	69.2	70.7	72.2	73.7	75.2	76.7	78.2	79.8	81.3	82.8	84.4	85.9	87.5	89.1	90.7	92.2	93.8
21	61.3	62.8	64.3	65.8	67.3	68.8	70.3	71.8	73.3	74.9	76.4	77.9	79.5	81.1	82.6	84.2	85.8	87.4	89.0	90.6	92.2	93.8	95.4	97.1
22	63.8	65.3	66.8	68.3	69.8	71.4	72.9	74.5	76.1	77.6	79.2	80.8	82.4	84.0	85.6	87.2	88.9	90.5	92.1	93.8	95.5	97.1	98.8	100.5
23	66.3	67.8	69.4	71.0	72.5	74.1	75.7	77.3	78.9	80.5	82.2	83.8	85.4	87.1	88.7	90.4	92.1	93.8	95.5	97.2	98.9	100.6	102.4	104.1
24	68.9	70.5	72.1	73.7	75.3	77.0	78.6	80.3	81.9	83.6	85.2	86.9	88.6	90.3	92.0	93.8	95.5	97.2	99.0	100.7	102.5	104.3	106.1	107.9
25	71.7	73.3	75.0	76.6	78.3	80.0	81.7	83.3	85.1	86.8	88.5	90.2	92.0	93.7	95.5	97.3	99.1	100.9	102.7	104.5	106.3	108.2	110.0	111.9
26	74.6	76.3	78.0	79.7	81.4	83.1	84.8	86.6	88.4	90.1	91.9	93.7	95.5	97.3	99.2	101.0	102.9	104.7	106.6	108.5	110.4	112.3	114.2	116.2
27	77.6	79.4	81.1	82.9	84.6	86.4	88.2	90.0	91.9	93.7	95.5	97.4	99.3	101.2	103.1	105.0	106.9	108.8	110.8	112.7	114.7	116.7	118.7	120.7
28	80.8	82.6	84.4	86.3	88.1	89.9	91.8	93.7	95.6	97.5	99.4	101.3	103.3	105.2	107.2	109.2	111.2	113.2	115.2	117.3	119.3	121.4	123.5	125.6
29	84.2	86.1	87.9	89.8	91.7	93.7	95.6	97.5	99.5	101.5	103.5	105.5	107.5	109.5	111.6	113.7	115.7	117.8	120.0	122.1	124.2	126.4	128.6	130.8
30	87.8	89.7	91.7	93.6	95.6	97.6	99.6	101.6	103.7	105.7	107.8	109.9	112.0	114.2	116.3	118.5	120.6	122.8	125.1	127.3	129.5	131.8	134.1	136.4
31	91.6	93.6	95.6	97.7	99.7	101.8	103.9	106.0	108.2	110.3	112.5	114.7	116.9	119.1	121.4	123.6	125.9	128.2	130.5	132.9	135.3	137.7	140.1	142.5
32	95.7	97.8	99.9	102.0	104.2	106.3	108.5	110.7	113.0	115.2	117.5	119.8	122.1	124.5	126.8	129.2	131.6	134.0	136.5	139.0	141.5	144.0	146.6	149.1
33	100.0	102.2	104.4	106.6	108.9	111.2	113.5	115.8	118.2	120.5	122.9	125.4	127.8	130.3	132.8	135.3	137.8	140.4	143.0	145.6	148.3	150.9	153.7	156.4
34	104.7	107.0	109.3	111.7	114.0	116.4	118.9	121.3	123.8	126.3	128.8	131.4	134.0	136.6	139.2	141.9	144.6	147.4	150.1	152.9	155.7	158.6	161.5	164.4
35	109.7	112.2	114.6	117.1	119.6	122.2	124.7	127.3	129.9	132.6	135.3	138.0	140.8	143.6	146.4	149.2	152.1	155.0	158.0	161.0	164.0	167.1	170.2	173.3
36	115.2	117.8	120.4	123.0	125.7	128.4	131.1	133.9	136.7	139.5	142.4	145.3	148.3	151.3	154.3	157.3	160.5	163.6	166.8	170.0	173.3	176.6	179.9	183.3
37	121.3	124.0	126.8	129.6	132.4	135.3	138.2	141.2	144.2	147.3	150.3	153.5	156.7	159.9	163.1	166.5	169.8	173.2	176.7	180.2	183.7	187.3	191.0	194.7
38	127.9	130.8	133.8	136.8	139.9	143.0	146.2	149.4	152.6	155.9	159.2	162.6	166.1	169.6	173.2	176.8	180.4	184.2	188.0	191.8	195.7	199.7	203.7	207.7
39	135.3	138.5	141.7	145.0	148.3	151.7	155.1	158.6	162.1	165.7	169.4	173.1	176.9	180.7	184.7	188.7	192.7	196.8	201.0	205.3	209.6	214.0	218.5	223.0
40	143.7	147.1	150.6	154.2	157.8	161.5	165.3	169.1	173.0	177.0	181.1	185.2	189.4	193.7	198.1	202.5	207.1	211.7	216.4	221.1	226.0	231.0	236.0	241.1
41	153.2	157.0	160.9	164.8	168.9	173.0	177.2	181.5	185.8	190.3	194.8	199.5	204.2	209.1	214.0	219.1	224.2	229.4	234.8	240.2	245.8	251.5	257.2	263.1
42	164.3	168.6	172.9	177.3	181.9	186.5	191.3	196.1	201.1	206.2	211.4	216.7	222.2	227.7	233.4	239.2	245.2	251.3	257.5	263.8	270.3	276.9	283.6	290.5
43	177.5	182.3	187.3	192.4	197.6	202.9	208.4	214.0	219.8	225.8	231.8	238.1	244.5	251.0	257.7	264.6	271.7	278.9	286.3	293.8	301.5	309.4	317.4	325.7
44	193.6	199.3	205.1	211.0	217.2	223.5	230.0	236.7	243.6	250.8	258.1	265.6	273.3	281.2	289.4	297.8	306.3	315.1	324.1	333.3	342.8	352.4	362.3	372.4
45	214.1																							

## Appendix B: Bacteria Sample Collection Form

Colilert (E. coli results)

Personnel: \_\_\_\_\_

[illegible]